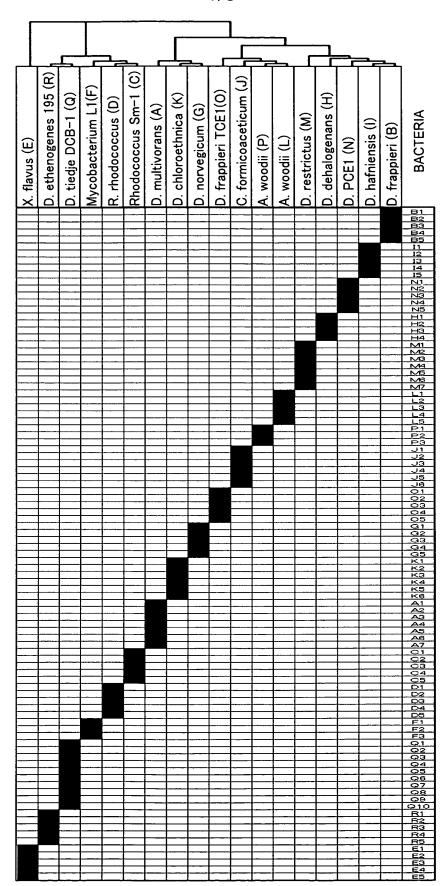
ITS SEQUENCE ALIGNMENT



PROBES

FIG. 1

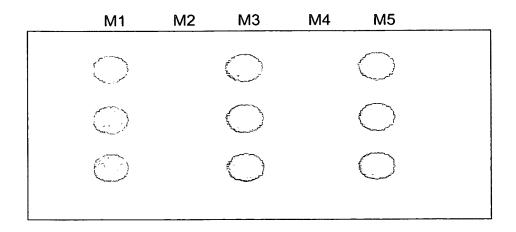


FIG. 2A

Contaminated soil sample hyrbidization

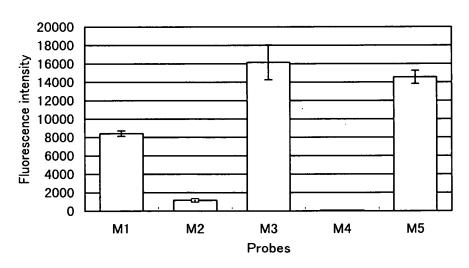


FIG. 2B

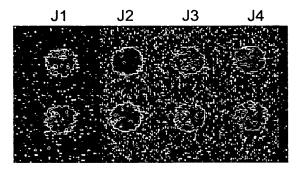


FIG. 3A

Contaminated groundwater sample hybridization

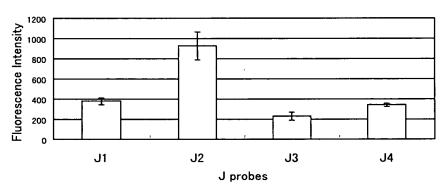
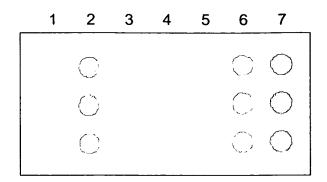
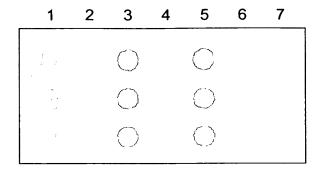


FIG. 3B

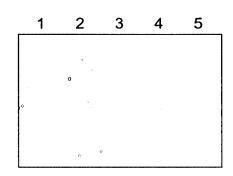
A probes (for Dehalospirillium multivorans)



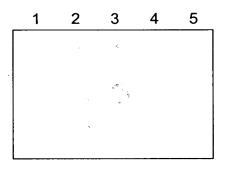
M probes (for Dehalobacter restrictus)



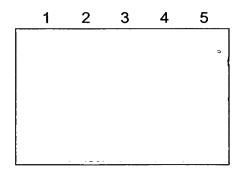
B probes (for *Desulfitobacterium frappieri*)



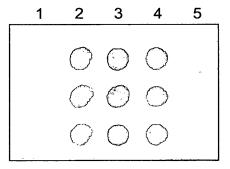
N probes (for *Desulfitobacterium* PCE1)



I probes (for *Desulfitobacterium hafniense*)



O probes (for Desulfitobacterium frappieri TCE1)



J probes (for Clostridium formicoaceticum)

1	2	3	4
0			

FIG. 5A

Dehalococcoides ethenogenes 195 R	$PCE \rightarrow TCE \rightarrow DCE \rightarrow VC \rightarrow ethene$
Desulfitobacterium frappieri B	PCE → TCE→ cisDCE
Desulfitobacterium hafniense I	
Desulfitobacterium dehalogenans H	
Desulfitobacterium sp. strain PCE1 N	
Desulfitobacterium frappieri TCE1 O	
Desulfomonile tiedjei DCB-1 Q	
Desulfuromonas chloroethenica K	PCE → TCE→ DCE
Acetobacterium woodii L	PCE → TCE
Acetobacterium woodii P	
Clostridium formicoaceticum J	PCE → TCE
Dehalobacter restrictus M	PCE → cisDCE
Dehalospirillum multivorans A	PCE → cisDCE
Desulfomicrobium norvegicum G	PCE → cisDCE
Rhodococcus sp. Sm-1 C	DEC, VC → CO2
Rhodococcus rhodococcus D	
<u> </u>	205 1/0 200
Xanthobacter flavus E	DGE, VG → CO2
	<u> </u>
Mycobacterium L1 F	VC → CO2

FIG. 5B